In the Claims

1 (currently amended). A method [of modulating] <u>for inhibiting</u> the function of <u>a</u> transcription [factors by] <u>factor</u>, <u>said method comprising</u> administering an effective amount of [an] <u>a</u> <u>double-stranded</u> oligonucleotide [containing optimal nucleotide binding sites for the transcription factor], <u>said oligonucleotide having a nucleotide sequence comprising the sequence TTCNNNGAA</u>, <u>wherein N represents any nucleotide</u>, and <u>wherein said transcription factor binds to said oligonucleotide</u>.

Claims 2-7 (canceled)

8 (currently amended). A [pharmaceutical] composition for inhibiting a transcription factor in a cell comprising [an effective amount of] a [double stranded] <u>double-stranded</u> oligonucleotide, said oligonucleotide having [a sequence bound by a transcription factor] <u>a nucleotide sequence comprising the sequence TTCNNNGAA</u>, wherein N represents any nucleotide, and wherein said transcription factor binds to said oligonucleotide.

9 (currently amended). The [pharmaceutical] composition according to claim [9] 8, wherein [in which] said transcription factor is activated.

10 (currently amended). The [pharmaceutical] composition according to claim 9, wherein said transcription factor is constitutively activated.

11 (currently amended). The [pharmaceutical] composition according to claim [9] 8, wherein the cell is a malignant cell.

12 (currently amended). The [pharmaceutical] composition according to claim [9] 8, wherein the cell is a leukemia cell.

13 (currently amended). The [pharmaceutical] composition according to claim 8, wherein said transcription factor is STAT5 [and said oligonucleotide contains the sequence TTCNNNGAA, in which "N" is any nucleotide].

Claim 14 (canceled)

15 (currently amended). The [pharmaceutical] composition according to claim 13, wherein said oligonucleotide [is selected from the group comprising an oligonucleotides having] comprises the sequence AGATTTCTAGGAATTCAAATC (SEQ ID NO:1)[, GCCTGATTTCCCCGAAATGACGGCA (SEQ ID NO:2) and GTATTTCCCAGAAAAGGAAC (SEQ ID NO:3)].

16 (currently amended). A method of inhibiting [malignant] proliferation of a tumor cell by administering an effective amount of a [double stranded] double-stranded oligonucleotide, [the] said oligonucleotide having [a sequence bound by a transcription factor,] a nucleotide sequence comprising the sequence TTCNNNGAA, wherein N represents any nucleotide, and wherein a transcription factor in said tumor cell binds to said oligonucleotide, the transcription factor activity being correlated to [malignant] proliferation of said tumor cell.

Claims 17-18 (canceled)

19 (currently amended). A method of removing [malignant] <u>a tumor</u> cell in vitro by exposing a cell culture to an effective amount of <u>a double-stranded</u> oligonucleotide [containing optimal nucleotide binding sites for a transcription factor], <u>said oligonucleotide having a nucleotide sequence comprising the sequence TTCNNNGAA</u>, wherein N represents any nucleotide, and wherein a <u>transcription factor in said tumor cell binds to said oligonucleotide</u>, the transcription factor activity being correlated to proliferation of said tumor cell.

20 (currently amended). [A therapeutic] <u>An</u> agent comprising an effective amount of [an] <u>a</u> double-stranded oligonucleotide [for modulating the function of transcription factors] <u>of claim 8</u> and a pharmaceutically effective carrier.

- 21 (new). The agent according to claim 20, wherein said oligonucleotide comprises the sequence AGATTCTAGGAATTCAAATC (SEQ ID NO:1).
 - 22 (new). The agent according to claim 20, wherein said transcription factor is STAT5.

- 23 (new). The agent according to claim 20, wherein said transcription factor is activated.
- 24 (new). The agent according to claim 23, wherein said transcription factor is constitutively activated.
 - 25 (new). The agent according to claim 20, wherein said cell is a malignant cell.
 - 26 (new). The agent according to claim 20, wherein said cell is a leukemia cell.
- 27 (new). The agent according to claim 20, wherein said oligonucleotide comprises multiple copies of said nucleotide sequence TTCNNNGAA.
- 28 (new). The agent according to claim 20, wherein said oligonucleotide comprises two copies of said nucleotide sequence TTCNNNGAA.
 - 29 (new). The agent according to claim 20, wherein said cell is a human cell.
 - 30 (new). The method according to claim 1, wherein said transcription factor is STAT5.
- 31 (new). The method according to claim 1, wherein said oligonucleotide comprises the sequence AGATTCTAGGAATTCAAATC (SEQ ID NO:1).
 - 32 (new). The method according to claim 1, wherein said transcription factor is activated.
- 33 (new). The method according to claim 32, wherein said transcription factor is constitutively activated.
- 34 (new). The method according to claim 1, wherein said oligonucleotide comprises multiple copies of said nucleotide sequence TTCNNNGAA.

- 35 (new). The method according to claim 1, wherein said oligonucleotide comprises two copies of said nucleotide sequence TTCNNNGAA.
- 36 (new). The composition according to claim 8, wherein said oligonucleotide comprises the sequence AGATTCTAGGAATTCAAATC (SEQ ID NO:1).
- 37 (new). The composition according to claim 8, wherein said oligonucleotide comprises multiple copies of said nucleotide sequence TTCNNNGAA.
- 38 (new). The composition according to claim 8, wherein said oligonucleotide comprises two copies of said nucleotide sequence TTCNNNGAA.
 - 39 (new). The composition according to claim 8, wherein said cell is a human cell.
- 40 (new). The method according to claim 16, wherein said oligonucleotide comprises the sequence AGATTCTAGGAATTCAAATC (SEQ ID NO:1).
 - 41 (new). The method according to claim 16, wherein said transcription factor is STAT5.
 - 42 (new). The method according to claim 16, wherein said transcription factor is activated.
- 43 (new). The method according to claim 42, wherein said transcription factor is constitutively activated.
 - 44 (new). The method according to claim 16, wherein said cell is a malignant cell.
 - 45 (new). The method according to claim 16, wherein said cell is a leukemia cell.

- 46 (new). The method according to claim 16, wherein said oligonucleotide comprises multiple copies of said nucleotide sequence TTCNNNGAA.
- 47 (new). The method according to claim 16, wherein said oligonucleotide comprises two copies of said nucleotide sequence TTCNNNGAA.
 - 48 (new). The method according to claim 16, wherein said cell is a human cell.
- 49 (new). The method according to claim 19, wherein said oligonucleotide comprises the sequence AGATTCTAGGAATTCAAATC (SEQ ID NO:1).
 - 50 (new). The method according to claim 19, wherein said transcription factor is STAT5.
 - 51 (new). The method according to claim 19, wherein said transcription factor is activated.
- 52 (new). The method according to claim 51, wherein said transcription factor is constitutively activated.
 - 53 (new). The method according to claim 19, wherein said cell is a malignant cell.
 - 54 (new). The method according to claim 19, wherein said cell is a leukemia cell.
- 55 (new). The method according to claim 19, wherein said oligonucleotide comprises multiple copies of said nucleotide sequence TTCNNNGAA.
- 56 (new). The method according to claim 19, wherein said oligonucleotide comprises two copies of said nucleotide sequence TTCNNNGAA.
 - 57 (new). The method according to claim 19, wherein said cell is a human cell.